

## **RAW SEQUENCE LISTING**

**The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.**

Application Serial Number: 10/542,704  
Source: PCT  
Date Processed by STIC: 07/28/2005

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PCT

## RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/542,704

DATE: 07/28/2005

TIME: 15:38:36

Input Set : A:\sequence listing ascii.txt

Output Set: N:\CRF4\07282005\J542704.raw

3 <110> APPLICANT: Gardella, Thomas J.  
 4 Kronenberg, Henry M.  
 5 Potts, John T.

7 &lt;120&gt; TITLE OF INVENTION: Conformationally Constrained Parathyroid Hormone (PTH)

## Analogs

8 With Lactam Bridges  
 10 <130> FILE REFERENCE: 0609.514PC00  
 C--> 12 <140> CURRENT APPLICATION NUMBER: US/10/542,704  
 C--> 12 <141> CURRENT FILING DATE: 2005-07-20  
 12 <160> NUMBER OF SEQ ID NOS: 37  
 14 <170> SOFTWARE: PatentIn version 3.2  
 16 <210> SEQ ID NO: 1  
 17 <211> LENGTH: 14  
 18 <212> TYPE: PRT  
 19 <213> ORGANISM: Artificial Sequence  
 21 <220> FEATURE:  
 22 <223> OTHER INFORMATION: Mutated hPTH  
 24 <220> FEATURE:  
 25 <221> NAME/KEY: MISC\_FEATURE  
 26 <222> LOCATION: (1)..(1)  
 27 <223> OTHER INFORMATION: Xaa can represent Gly, Ser, Ala or alpha-aminoisobutyric acid  
 29 <220> FEATURE:  
 30 <221> NAME/KEY: MISC\_FEATURE  
 31 <222> LOCATION: (3)..(3)  
 32 <223> OTHER INFORMATION: Xaa can represent Ala, Ser, alpha-aminoisobutyric acid  
 34 <220> FEATURE:  
 35 <221> NAME/KEY: MISC\_FEATURE  
 36 <222> LOCATION: (6)..(6)  
 37 <223> OTHER INFORMATION: Xaa can represent Asp, Glu or Lys  
 39 <220> FEATURE:  
 40 <221> NAME/KEY: MISC\_FEATURE  
 41 <222> LOCATION: (10)..(10)  
 42 <223> OTHER INFORMATION: Xaa can represent Asp, Glu or Lys  
 44 <220> FEATURE:  
 45 <221> NAME/KEY: MISC\_FEATURE  
 46 <222> LOCATION: (11)..(11)  
 47 <223> OTHER INFORMATION: Xaa can represent Arg, Leu or Homoarginine  
 49 <220> FEATURE:  
 50 <221> NAME/KEY: MISC\_FEATURE  
 51 <222> LOCATION: (12)..(12)  
 52 <223> OTHER INFORMATION: Xaa can represent Ala or Gly  
 54 <220> FEATURE:  
 55 <221> NAME/KEY: MISC\_FEATURE  
 56 <222> LOCATION: (14)..(14)

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RAW SEQUENCE LISTING DATE: 07/28/2005  
 PATENT APPLICATION: US/10/542,704 TIME: 15:38:36

Input Set : A:\sequence listing ascii.txt  
 Output Set: N:\CRF4\07282005\J542704.raw

57 <223> OTHER INFORMATION: Xaa can represent Trp or His  
 59 <220> FEATURE:  
 60 <221> NAME/KEY: MOD\_RES  
 61 <222> LOCATION: (14)..(14)  
 62 <223> OTHER INFORMATION: AMIDATION  
 64 <400> SEQUENCE: 1  
 W--> 66 Xaa Val Xaa Glu Ile Xaa Leu Met His Xaa Xaa Xaa Lys Xaa  
 67 1 5 10  
 70 <210> SEQ ID NO: 2  
 71 <211> LENGTH: 12  
 72 <212> TYPE: PRT  
 73 <213> ORGANISM: Artificial Sequence  
 75 <220> FEATURE:  
 76 <223> OTHER INFORMATION: Mutated hPTH  
 79 <220> FEATURE:  
 80 <221> NAME/KEY: MISC\_FEATURE  
 81 <222> LOCATION: (6)..(6)  
 82 <223> OTHER INFORMATION: Xaa can represent Asp, Glu or Lys  
 84 <220> FEATURE:  
 85 <221> NAME/KEY: MISC\_FEATURE  
 86 <222> LOCATION: (9)..(9)  
 87 <223> OTHER INFORMATION: Xaa represents homoarginine  
 89 <220> FEATURE:  
 90 <221> NAME/KEY: MOD\_RES  
 91 <222> LOCATION: (12)..(12)  
 92 <223> OTHER INFORMATION: AMIDATION  
 94 <400> SEQUENCE: 2  
 W--> 96 Ala Val Ala Glu Ile Xaa Leu Met Xaa Ala Lys Trp  
 97 1 5 10  
 100 <210> SEQ ID NO: 3  
 101 <211> LENGTH: 14  
 102 <212> TYPE: PRT  
 103 <213> ORGANISM: Artificial Sequence  
 105 <220> FEATURE:  
 106 <223> OTHER INFORMATION: [(Glu-6, Lys-10) lac.] PTH(1-14)  
 109 <220> FEATURE:  
 110 <221> NAME/KEY: MISC\_FEATURE  
 111 <222> LOCATION: (1)..(14)  
 112 <223> OTHER INFORMATION: Sequence is cyclized via a 6-10 lactam bridge  
 114 <220> FEATURE:  
 115 <221> NAME/KEY: MISC\_FEATURE  
 116 <222> LOCATION: (11)..(11)  
 117 <223> OTHER INFORMATION: Xaa represents homoarginine  
 119 <220> FEATURE:  
 120 <221> NAME/KEY: MOD\_RES  
 121 <222> LOCATION: (14)..(14)  
 122 <223> OTHER INFORMATION: AMIDATION  
 124 <400> SEQUENCE: 3  
 W--> 126 Ala Val Ala Glu Ile Glu Leu Met His Lys Xaa Ala Lys Trp

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PATENT APPLICATION: US/10/542,704

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TIME: 15:38:36

Input Set : A:\sequence listing ascii.txt  
 Output Set: N:\CRF4\07282005\J542704.raw

127 1 5 10  
 130 <210> SEQ ID NO: 4  
 131 <211> LENGTH: 14  
 132 <212> TYPE: PRT  
 133 <213> ORGANISM: Artificial Sequence  
 135 <220> FEATURE:  
 136 <223> OTHER INFORMATION: [(Lys-6, Lys-10) lac.] PTH(1-14)  
 139 <220> FEATURE:  
 140 <221> NAME/KEY: MISC\_FEATURE  
 141 <222> LOCATION: (1)..(14)  
 142 <223> OTHER INFORMATION: Sequence is cyclized via 6-10 lactam bridge  
 144 <220> FEATURE:  
 145 <221> NAME/KEY: MOD\_RES  
 146 <222> LOCATION: (14)..(14)  
 147 <223> OTHER INFORMATION: AMIDATION  
 149 <400> SEQUENCE: 4  
 W--> 151 Ala Val Ala Glu Ile Lys Leu Met His Lys Xaa Ala Lys Trp  
 152 1 5 10  
 155 <210> SEQ ID NO: 5  
 156 <211> LENGTH: 34  
 157 <212> TYPE: PRT  
 158 <213> ORGANISM: Rattus sp.  
 161 <220> FEATURE:  
 162 <221> NAME/KEY: MOD\_RES  
 163 <222> LOCATION: (34)..(34)  
 164 <223> OTHER INFORMATION: AMIDATION  
 166 <400> SEQUENCE: 5  
 168 Ala Val Ser Glu Ile Gln Leu Met His Asn Leu Gly Lys His Leu Ala  
 169 1 5 10 15  
 172 Ser Val Glu Arg Met Gln Trp Leu Arg Lys Lys Leu Gln Asp Val His  
 173 20 25 30  
 176 Asn Phe  
 180 <210> SEQ ID NO: 6  
 181 <211> LENGTH: 34  
 182 <212> TYPE: PRT  
 183 <213> ORGANISM: Homo sapiens  
 186 <220> FEATURE:  
 187 <221> NAME/KEY: MOD\_RES  
 188 <222> LOCATION: (34)..(34)  
 189 <223> OTHER INFORMATION: AMIDATION  
 191 <400> SEQUENCE: 6  
 193 Ser Val Ser Glu Ile Gln Leu Met His Asn Leu Gly Lys His Leu Asn  
 194 1 5 10 15  
 197 Ser Met Glu Arg Val Glu Trp Leu Arg Lys Lys Leu Gln Asp Val His  
 198 20 25 30  
 201 Asn Phe  
 205 <210> SEQ ID NO: 7  
 206 <211> LENGTH: 34  
 207 <212> TYPE: PRT

## RAW SEQUENCE LISTING

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TIME: 15:38:36

Input Set : A:\sequence listing ascii.txt  
Output Set: N:\CRF4\07282005\J542704.raw

208 <213> ORGANISM: Artificial Sequence  
210 <220> FEATURE:  
211 <223> OTHER INFORMATION: Mutated hPTH  
214 <220> FEATURE:  
215 <221> NAME/KEY: MISC\_FEATURE  
216 <222> LOCATION: (6)..(6)  
217 <223> OTHER INFORMATION: Xaa can represent Ala, Glu or Gln  
219 <220> FEATURE:  
220 <221> NAME/KEY: MISC\_FEATURE  
221 <222> LOCATION: (10)..(10)  
222 <223> OTHER INFORMATION: Xaa can represent Ala, Glu or Gln  
224 <220> FEATURE:  
225 <221> NAME/KEY: MOD\_RES  
226 <222> LOCATION: (34)..(34)  
227 <223> OTHER INFORMATION: AMIDATION  
229 <400> SEQUENCE: 7  
W--> 231 Ser Val Ser Glu Ile Xaa Leu Met His Xaa Leu Gly Lys His Leu Asn  
232 1 5 10 15  
235 Ser Met Glu Arg Val Glu Trp Leu Arg Lys Lys Leu Gln Asp Val His  
236 20 25 30  
239 Asn Tyr  
243 <210> SEQ ID NO: 8  
244 <211> LENGTH: 11  
245 <212> TYPE: PRT  
246 <213> ORGANISM: Artificial Sequence  
248 <220> FEATURE:  
249 <223> OTHER INFORMATION: Mutated hPTH(1-11)  
252 <220> FEATURE:  
253 <221> NAME/KEY: MOD\_RES  
254 <222> LOCATION: (11)..(11)  
255 <223> OTHER INFORMATION: AMIDATION  
257 <400> SEQUENCE: 8  
259 Ala Val Ala Glu Ile Gln Leu Met His Gln Arg  
260 1 5 10  
263 <210> SEQ ID NO: 9  
264 <211> LENGTH: 14  
265 <212> TYPE: PRT  
266 <213> ORGANISM: Artificial Sequence  
268 <220> FEATURE:  
269 <223> OTHER INFORMATION: [Aib-1,3]PTH(1-14)  
272 <220> FEATURE:  
273 <221> NAME/KEY: MISC\_FEATURE  
274 <222> LOCATION: (1)..(1)  
275 <223> OTHER INFORMATION: Xaa represents alpha-aminoisobutyric acid  
277 <220> FEATURE:  
278 <221> NAME/KEY: MISC\_FEATURE  
279 <222> LOCATION: (3)..(3)  
280 <223> OTHER INFORMATION: Xaa represents alpha-aminoisobutyric acid  
282 <220> FEATURE:

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TIME: 15:38:36

Input Set : A:\sequence listing ascii.txt  
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283 <221> NAME/KEY: MISC\_FEATURE  
284 <222> LOCATION: (11)..(11)  
285 <223> OTHER INFORMATION: Xaa represents homoarginine  
287 <220> FEATURE:  
288 <221> NAME/KEY: MOD\_RES  
289 <222> LOCATION: (14)..(14)  
290 <223> OTHER INFORMATION: AMIDATION  
292 <400> SEQUENCE: 9  
W--> 294 Xaa Val Xaa Glu Ile Gln Leu Met His Gln Xaa Ala Lys Trp  
295 1 5 10  
298 <210> SEQ ID NO: 10  
299 <211> LENGTH: 34  
300 <212> TYPE: PRT  
301 <213> ORGANISM: Artificial Sequence  
303 <220> FEATURE:  
304 <223> OTHER INFORMATION: ([Tyr-34]hPTH(1-34)amide)  
307 <220> FEATURE:  
308 <221> NAME/KEY: MOD\_RES  
309 <222> LOCATION: (34)..(34)  
310 <223> OTHER INFORMATION: AMIDATION  
312 <400> SEQUENCE: 10  
314 Ser Val Ser Glu Ile Gln Leu Met His Asn Leu Gly Lys His Leu Asn  
315 1 5 10 15  
318 Ser Met Glu Arg Val Glu Trp Leu Arg Lys Lys Leu Gln Asp Val His  
319 20 25 30  
322 Asn Tyr  
326 <210> SEQ ID NO: 11  
327 <211> LENGTH: 14  
328 <212> TYPE: PRT  
329 <213> ORGANISM: Artificial Sequence  
331 <220> FEATURE:  
332 <223> OTHER INFORMATION: [Ala-1,3,12, Gln-10, Har-11, Trp-14]PTH(1-14)amide  
335 <220> FEATURE:  
336 <221> NAME/KEY: MISC\_FEATURE  
337 <222> LOCATION: (11)..(11)  
338 <223> OTHER INFORMATION: Xaa represents homoarginine  
340 <220> FEATURE:  
341 <221> NAME/KEY: MOD\_RES  
342 <222> LOCATION: (14)..(14)  
343 <223> OTHER INFORMATION: AMIDATION  
345 <400> SEQUENCE: 11  
W--> 347 Ala Val Ala Glu Ile Gln Leu Met His Gln Xaa Ala Lys Trp  
348 1 5 10  
351 <210> SEQ ID NO: 12  
352 <211> LENGTH: 11  
353 <212> TYPE: PRT  
354 <213> ORGANISM: Artificial Sequence  
356 <220> FEATURE:  
357 <223> OTHER INFORMATION: [Ala-1,3, Gln-10, Har-11]PTH(1-11)amide

RAW SEQUENCE LISTING ERROR SUMMARY  
PATENT APPLICATION: US/10/542,704

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TIME: 15:38:37

Input Set : A:\sequence listing ascii.txt  
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Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:1; Xaa Pos. 1,3,6,10,11,12,14  
Seq#:2; Xaa Pos. 6,9  
Seq#:3; Xaa Pos. 11  
Seq#:4; Xaa Pos. 11  
Seq#:7; Xaa Pos. 6,10  
Seq#:9; Xaa Pos. 1,3,11  
Seq#:11; Xaa Pos. 11  
Seq#:12; Xaa Pos. 11  
Seq#:14; Xaa Pos. 11  
Seq#:15; Xaa Pos. 1,3,11  
Seq#:16; Xaa Pos. 11  
Seq#:17; Xaa Pos. 1,3,11  
Seq#:21; Xaa Pos. 1,3,11  
Seq#:22; Xaa Pos. 1,3,11  
Seq#:23; Xaa Pos. 1,3,11  
Seq#:24; Xaa Pos. 11  
Seq#:25; Xaa Pos. 11  
Seq#:26; Xaa Pos. 11  
Seq#:30; Xaa Pos. 11  
Seq#:31; Xaa Pos. 1,3,6,10,11,12,14  
Seq#:32; Xaa Pos. 1,3,6,10,11,12,14  
Seq#:33; Xaa Pos. 1,3,6,10,11,12  
Seq#:34; Xaa Pos. 1,3,6,10,11,12  
Seq#:35; Xaa Pos. 1,3,6,10,11

## VERIFICATION SUMMARY

PATENT APPLICATION: US/10/542,704

DATE: 07/28/2005

TIME: 15:38:37

Input Set : A:\sequence listing ascii.txt  
Output Set: N:\CRF4\07282005\J542704.raw

L:12 M:270 C: Current Application Number differs, Replaced Current Application No  
L:12 M:271 C: Current Filing Date differs, Replaced Current Filing Date  
L:66 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:0  
L:96 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:0  
L:126 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3 after pos.:0  
L:151 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4 after pos.:0  
L:231 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7 after pos.:0  
L:294 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9 after pos.:0  
L:347 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:11 after pos.:0  
L:372 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:12 after pos.:0  
L:434 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:14 after pos.:0  
L:473 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:15 after pos.:0  
L:502 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:16 after pos.:0  
L:541 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:17 after pos.:0  
L:640 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:21 after pos.:0  
L:675 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:22 after pos.:0  
L:710 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:23 after pos.:0  
L:735 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:24 after pos.:0  
L:760 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:25 after pos.:0  
L:790 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:26 after pos.:0  
L:900 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:30 after pos.:0  
L:959 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:31 after pos.:0  
L:1022 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:32 after pos.:0  
L:1076 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:33 after pos.:0  
L:1126 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:34 after pos.:0  
L:1171 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:35 after pos.:0